

Hazar - Wesley

RECEIVED 646

MAY 10 2001

TECH CENTER 1600/2900

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/383,745

DATE: 05/04/2001

TIME: 12:39:12

Input Set : N:\Crf3\RULE60\09383745.txt

Output Set: N:\CRF3\05042001\I383745.raw

ENTERED

3 <110> APPLICANT: Glucksmann, Maria A.
5 <120> TITLE OF INVENTION: 14926 Receptor, A Novel G-Protein Coupled Receptor
7 <130> FILE REFERENCE: 035800/169197
9 <140> CURRENT APPLICATION NUMBER: 09/383,745
10 <141> CURRENT FILING DATE: 1999-08-26
12 <150> PRIOR APPLICATION NUMBER: 09/145,745
13 <151> PRIOR FILING DATE: 1998-09-02
15 <160> NUMBER OF SEQ ID NOS: 3
17 <170> SOFTWARE: PatentIn Ver. 2.1
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 370
21 <212> TYPE: PRT
22 <213> ORGANISM: Homo sapiens
24 <400> SEQUENCE: 1
25 Met Ala Asn Tyr Ser His Ala Ala Asp Asn Ile Leu Gln Asn Leu Ser
26 1 5 10 15
28 Pro Leu Thr Ala Phe Leu Lys Leu Thr Ser Leu Gly Phe Ile Ile Gly
29 20 25 30
31 Val Ser Val Val Gly Asn Leu Leu Ile Ser Ile Leu Leu Val Lys Asp
32 35 40 45
34 Lys Thr Leu His Arg Ala Pro Tyr Tyr Phe Leu Leu Asp Leu Cys Cys
35 50 55 60
37 Ser Asp Ile Leu Arg Ser Ala Ile Cys Phe Pro Phe Val Phe Asn Ser
38 65 70 75 80
40 Val Lys Asn Gly Ser Thr Trp Thr Tyr Gly Thr Leu Thr Cys Lys Val
41 85 90 95
43 Ile Ala Phe Leu Gly Val Leu Ser Cys Phe His Thr Ala Phe Met Leu
44 100 105 110
46 Phe Cys Ile Ser Val Thr Arg Tyr Leu Ala Ile Ala His Arg Phe
47 115 120 125
49 Tyr Thr Lys Arg Leu Thr Phe Trp Thr Cys Leu Ala Val Ile Cys Met
50 130 135 140
52 Val Trp Thr Leu Ser Val Ala Met Ala Phe Pro Pro Val Leu Asp Val
53 145 150 155 160
55 Gly Thr Tyr Ser Phe Ile Arg Glu Glu Asp Gln Cys Thr Phe Gln His
56 165 170 175
58 Arg Ser Phe Arg Ala Asn Asp Ser Leu Gly Phe Met Leu Leu Leu Ala
59 180 185 190
61 Leu Ile Leu Leu Ala Thr Gln Leu Val Tyr Leu Lys Leu Ile Phe Phe
62 195 200 205
64 Val His Asp Arg Arg Lys Met Lys Pro Val Gln Phe Val Ala Ala Val
65 210 215 220
67 Ser Gln Asn Trp Thr Phe His Gly Pro Gly Ala Ser Gly Gln Ala Ala
68 225 230 235 240
70 Ala Asn Trp Leu Ala Gly Phe Gly Arg Gly Pro Thr Pro Pro Thr Leu
71 245 250 255
73 Leu Gly Ile Arg Gln Asn Ala Asn Thr Thr Gly Arg Arg Arg Leu Leu